












Sublist	Category	Term	RT	Genes	Count	%	P-Value	Benjamini
<input type="checkbox"/>	KEGG_PATHWAY	Pathways in cancer	RT		20	3.6	3.3E-2	1.0E0
<input type="checkbox"/>	KEGG_PATHWAY	Cytokine-cytokine receptor interaction	RT		16	2.9	3.0E-3	4.4E-1
<input type="checkbox"/>	KEGG_PATHWAY	PI3K-Akt signaling pathway	RT		16	2.9	1.5E-2	8.6E-1
<input type="checkbox"/>	KEGG_PATHWAY	Axon guidance	RT		10	1.8	2.3E-2	1.0E0
<input type="checkbox"/>	KEGG_PATHWAY	Focal adhesion	RT		9	1.6	8.9E-2	1.0E0
<input type="checkbox"/>	KEGG_PATHWAY	ECM-receptor interaction	RT		8	1.4	3.8E-3	4.4E-1
<input type="checkbox"/>	KEGG_PATHWAY	Phagosome	RT		8	1.4	5.9E-2	1.0E0
<input type="checkbox"/>	KEGG_PATHWAY	Hippo signaling pathway	RT		8	1.4	6.7E-2	1.0E0
<input type="checkbox"/>	KEGG_PATHWAY	Protein digestion and absorption	RT		7	1.3	3.0E-2	1.0E0
<input type="checkbox"/>	KEGG_PATHWAY	Cholinergic synapse	RT		7	1.3	4.4E-2	1.0E0
<input type="checkbox"/>	KEGG_PATHWAY	Relaxin signaling pathway	RT		7	1.3	7.4E-2	1.0E0
<input type="checkbox"/>	KEGG_PATHWAY	Hedgehog signaling pathway	RT		6	1.1	8.7E-3	6.6E-1
<input type="checkbox"/>	KEGG_PATHWAY	Amoebiasis	RT		6	1.1	8.3E-2	1.0E0
<input type="checkbox"/>	KEGG_PATHWAY	Histidine metabolism	RT		3	0.5	8.9E-2	1.0E0